

# Sequence Listing

<110> Patricia Billing-Medel  
Maurice Cohen  
Tracey L. Colpitts  
Paula N. Friedman  
Julian Gordon  
Edward N. Granados  
Steven C. Hodges  
Michael R. Klass  
Jon D. Kratochvil  
Lisa Roberts-Rapp  
John C. Russell  
Stephen D. Stroupe

<120> Reagents and Methods Useful for Detecting Diseases of the Breast

<130> 6193.US.P1

<140> US 09/193,538

<141> 1998-11-17

<150> US 08/971,772

<151> 1997-11-17

<160> 23

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 288

<212> DNA

<213> Homo sapiens

<400> 1

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aggcctggag	accagctccg	gtgggaagct	ggctggccat	cagaagaccg	tccccacggc	180
tcacctgact	tttggttattg	actgcaccca	cgggaagcag	ctctccctgg	cagcaaccgc	240
atcaccaccc	caagccccca	gtcccaatcg	agggttgctc	ccccacca		288

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<211> 250

<212> DNA

<213> Homo sapiens

<400> 2

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gtgggaagct	ggctggccat	cagaagaccg	tccccacggc	tcacctgact	tttggttattg	120
actgcaccca	cgggaagcag	ctctccctgg	cagcaaccgc	atcaccaccc	caagccccca	180
gtcccaatcg	agggttctgc	acccaccaa	tgaagaccta	catcgtgttc	tgtggggaaa	240
actggccccca						250

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<211> 256

<212> DNA

<213> Homo sapiens

<400> 3

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cagcccgtc	tgccccccagg	aggttcccga	ggctaagggg	aaaccctga	aggctgcgcc	180
tgtgaggtct	tcaacttggg	gaacagtcaa	ggactcactg	aaagccctct	cctcttgtgt	240
ctgtgggcag	gccgat					256

<210> 4  
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ctccaggaaa accatggtat ctccccagca ctttgcaggg cctggcatgt ggaagatgta 120  
ccagtaatat ttgctgtatg aatgaatgag tctcttcatg tgcagggtgac ttatcctgcc 180  
tctgccactc gacggatggt tcagatgccc cttagcggat ctaatgatgt ttccttgggt 240  
caagcacaaa agactc 256

<210> 5  
<211> 133  
<212> DNA  
<213> Homo sapiens

<400> 5  
gctgttcaaa atcatcttct ttattttattg ggttacttta tttattcagg gtgggttccc 60  
tccaccccaa aaataaccagc tccaggaaaa ccatggtatc tccccagcac tttgcagggc 120  
ctggcatgtg gaa 133

<210> 6  
<211> 910  
<212> DNA  
<213> Homo sapiens

<400> 6  
agagtggcct aggacagctc ctctcctgcc agagctaggg aggcgcgcgaa gtagccgcat 60  
ggccccgtca gaagacccca gggactggag agccaacctc aaaggcacca tccgtgagac 120  
aggcctggag accagctccg gtgggaagct ggctggccat cagaagaccg tccccacggc 180  
tcacctgact tttgttattg actgcaccca cgggaagcag ctctccctgg cagcaaccgc 240  
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cttcccagtc agcccgtct gccccagga ggttcccag gctaaggagg aaccctgaa 480  
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aaggaaacat cattagatcc gctaaggggc atctgaaaca tccgtcgagt ggcagaggca 720  
ggataagtca cctgcacatg aagagactca ttcattcata cagcaaatat tactggtaca 780  
tcttccacat gccaggccct gcaaagtgtc ggggagatac catgggtttc ctggagctgg 840  
tatttttggg gtggagggaa cccaccctga ataaataaag taaccacaata aataaagaag 900  
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<212> DNA  
<213> Homo sapiens

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ggccccgtca gaagacccca gggactggag agccaacctc aaaggcacca tccgtgagac 120  
aggcctggag accagctccg gtgggaagct ggctggccat cagaagaccg tccccacggc 180  
tcacctgact tttgttattg actgcaccca cgggaagcag ctctccctgg cagcaaccgc 240  
atcaccaccc caagccccca gtcccaatcg agggcttgct accccaccaa tgaagacct 300  
catcgtgttc tgtggggaaa actggcccca tctkactcgg gtgaccccca tgggtggggg 360  
atgccttgcc caggccaggg ccaccctgcc gctctgcaga gggctctgtg cctcagcttc 420  
cttcccagtc agcccgtct gccccagga ggttcccag gctaaggagg aaccctgaa 480  
ggctgcgcct gtgaggtctt caacttgggg aacagtcaag gactcactga aagccctctc 540  
ctcttggtgc tgtgggcagg ccgattagct ggaagggccg ggctctgatg cccagaggct 600  
gcaattccca gggcctggcc ctgcttcccc agctaagcag gagtcttttg tgcttgagcc 660  
aaggaaacat cattagatcc gctaaggggc atctgaaaca tccgtcgagt ggcagaggca 720  
ggataagtca cctgcacatg aagagactca ttcattcata cagcaaatat tactggtaca 780  
tcttccacat gccaggccct gcaaagtgtc ggggagatac catgggtttc ctggagctgg 840  
tatttttggg gtggagggaa cccaccctga ataaataaag taaccacaata aataaagaag 900  
atgattttga acagc 915

<210> 8  
 <211> 68  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Restriction site  
  
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 cgggaatt 68  
  
 <210> 9  
 <211> 68  
 <212> DNA  
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 <223> Restriction site  
  
 <400> 9  
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 gaattccg 68  
  
 <210> 10  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Universal primer  
  
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 agcggataac aatttcacac agga 24  
  
 <210> 11  
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 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Universal primer  
  
 <400> 11  
 tgtaaaacga cggccagt 18  
  
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 <212> DNA  
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 <400> 12  
 cccacacat gaagacctac 20  
  
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 <213> Homo sapiens  
  
 <400> 13  
 agaggagagg gctttcagt 20  
  
 <210> 14  
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 <212> DNA  
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<400> 14  
ccccacagaa cacgatgtag

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<210> 15  
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<212> DNA  
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<400> 15  
ttgtcacccc accaatgaag ac

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<210> 16  
<211> 22  
<212> DNA  
<213> Homo sapiens

<400> 16  
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22

<210> 17  
<211> 188  
<212> PRT  
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<400> 17  
Glu Trp Pro Arg Thr Ala Pro Leu Leu Pro Glu Leu Gly Arg Arg Arg  
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Ser Ser Arg Met Ala Pro Ser Glu Asp Pro Arg Asp Trp Arg Ala Asn  
20 25 30  
Leu Lys Gly Thr Ile Arg Glu Thr Gly Leu Glu Thr Ser Ser Gly Gly  
35 40 45  
Lys Leu Ala Gly His Gln Lys Thr Val Pro Thr Ala His Leu Thr Phe  
50 55 60  
Val Ile Asp Cys Thr His Gly Lys Gln Leu Ser Leu Ala Ala Thr Ala  
65 70 75 80  
Ser Pro Pro Gln Ala Pro Ser Pro Asn Arg Gly Leu Val Thr Pro Pro  
85 90 95  
Met Lys Thr Tyr Ile Val Phe Cys Gly Glu Asn Trp Pro His Leu Thr  
100 105 110  
Arg Val Thr Pro Met Gly Gly Gly Cys Leu Ala Gln Ala Arg Ala Thr  
115 120 125  
Leu Pro Leu Cys Arg Gly Ser Val Ala Ser Ala Ser Phe Pro Val Ser  
130 135 140  
Pro Leu Cys Pro Gln Glu Val Pro Glu Ala Lys Gly Lys Pro Val Lys  
145 150 155 160  
Ala Ala Pro Val Arg Ser Ser Thr Trp Gly Thr Val Lys Asp Ser Leu  
165 170 175  
Lys Ala Leu Ser Ser Cys Val Cys Gly Gln Ala Asp  
180 185

<210> 18  
<211> 21  
<212> PRT  
<213> Homo sapiens

<400> 18  
Arg Ser Ser Arg Met Ala Pro Ser Glu Asp Pro Arg Asp Trp Arg Ala  
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Asn Leu Lys Gly Thr  
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<210> 19  
<211> 19  
<212> PRT  
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<400> 19  
Met Gly Gly Gly Cys Leu Ala Gln Ala Arg Ala Thr Leu Pro Leu Cys  
1 5 10 15  
Arg Gly Ser

<210> 20  
<211> 35  
<212> PRT  
<213> Homo sapiens

<400> 20  
Leu Cys Pro Gln Glu Val Pro Glu Ala Lys Gly Lys Pro Val Lys Ala  
1 5 10 15  
Ala Pro Val Arg Ser Ser Thr Trp Gly Thr Val Lys Asp Ser Leu Lys  
20 25 30  
Ala Leu Ser  
35

<210> 21  
<211> 19  
<212> PRT  
<213> Homo sapiens

<400> 21  
Arg Glu Thr Gly Leu Glu Thr Ser Ser Gly Gly Lys Leu Ala Gly His  
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Gln Lys Thr

<210> 22  
<211> 8  
<212> PRT  
<213> Artificial Sequence

<220>  
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<400> 22  
Asp Tyr Lys Asp Asp Asp Asp Lys  
1 5

<210> 23  
<211> 21  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Affinity purification system recognition site

<400> 23  
Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Met His Thr Glu His  
1 5 10 15  
His His His His His  
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